



1600

RAW SEQUENCE LISTING

PATENT APPLICATION: US/08/826,361B

DATE: 08/21/2003

TIME: 16:25:18

RECEIVED

AUG 28 2003

TECH CENTER 1600/2900

Input Set : A:\2355-124 Seq List.txt

Output Set: N:\CRF4\08212003\H826361B.raw

ENTERED

P. 6

3 <110> APPLICANT: Mosselman, Sietse
 4 Dijkema, E. in.
 5 <120> TITLE OF INVENTION: Novel Estrogen Receptor
 6 <130> FILE REFERENCE: 0/96193 US

C--> 10 <140> CURRENT APPLICATION NUMBER: US/08/826,361B

C--> 10 <141> CURRENT FILING DATE: 1997-03-26

10 <150> PRIOR APPLICATION NUMBER: US 08/826,361

11 <151> PRIOR FILING DATE: 1997-03-26

12 <150> PRIOR APPLICATION NUMBER: EP 96205294.3

13 <151> PRIOR FILING DATE: 1996-11-22

14 <150> PRIOR APPLICATION NUMBER: EP 96200820.7

15 <151> PRIOR FILING DATE: 1996-03-26

16 <160> NUMBER OF SEQ ID NOS: 32

17 <170> SOFTWARE: PatentIn version 3.0

18 <210> SEQ ID NO: 1

19 <211> LENGTH: 1434

20 <212> TYPE: DNA

21 <213> ORGANISM: Homo sapiens

22 <400> SEQUENCE: 1

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| 10 | atgaatttccg | gaatttccag | caatttccact | aacttgggaag | gtggggcctgg | tgggcagacc | 60 |
| 20 | aaagacccaa | atgtgtgtgtg | gccaacacat | gggacattt | ctcctttagt | ggtccatcgc | 120 |
| 30 | cagttatcac | atctgtgttg | gggaacotcaa | aagagtccct | ggtgtgaagc | aagatcgcga | 180 |
| 40 | gaacacacat | taccttcaa | cajajagaca | ctgaaaagga | aggttagtgg | gaacccgttgc | 240 |
| 50 | gccaagccctg | ttactgttcc | aggttcaaaag | agggatgctc | aacttcctggc | tgtctggcagc | 300 |
| 60 | gattacgcat | cgggatatca | ctatggagtc | tggctgtgtg | aagpatgtaa | ggcctttttt | 360 |
| 70 | aaagaagca | ttcaaaagaca | taattattat | atttgtccag | ctacaaatca | gtgtacaatc | 420 |
| 80 | gataaaacac | ggcgcaagag | ctgcccaggcc | tgcgcacttc | ggaagtgtta | cgaagtggga | 480 |
| 90 | atggtgaagt | gtggctcccg | cajajagaga | tgtgggtacc | gccttgtggc | gajacagaga | 540 |
| 100 | atgtccgaag | agcagttgca | ctgtgcctgg | aaggccaaga | gaagtggcgg | ccacgcgcgc | 600 |
| 110 | agagtggcgg | agctgtgtgt | cgaccccttg | agccccgagc | agctagtgtt | ccacctcctg | 660 |
| 120 | agaggtgagc | cgccctctgt | ccgtatcagc | cgccccagtg | cgcccttcac | cgaggccttc | 720 |
| 130 | atgatgatgt | ccctgaccaa | gttgcccgac | aaggagttag | tacacatgat | cagctggggc | 780 |
| 140 | aaagaagatt | ccggcttctg | ggaghtcagc | ctgttcgacc | aagtgcgggt | cttygagagc | 840 |
| 150 | tgttgatggt | aggtgttaat | galtggggctg | atgtgggcgt | caattgaaca | ccccgggaag | 900 |
| 160 | ctcatctttg | ctccagatct | tgttctggac | agggatgagg | ggaatgggt | agaagggaatt | 960 |
| 170 | ctggaaatct | ttgaacttgt | ccgtggaaact | aactcaaggt | ttcgagaggt | aaaactccaa | 1020 |
| 180 | caacaagaat | atctctgtgt | caaggccatg | atcctgtcca | attccagtat | gtacccctctg | 1080 |
| 190 | gtccacagcga | cccaggttgc | tgacagcagc | cggaagctgg | ctcaacttgt | gaacgccttg | 1140 |
| 200 | acccatgctt | tggtttttgt | galttccaaag | aggggcacat | cttcccagca | gcaatccatg | 1200 |
| 210 | cgcctggcta | acctcctgat | ccctctgttc | caagtcaggc | atgcgagtaa | caagggcctg | 1260 |
| 220 | gascctcttg | tcaacatgaa | gtgcacaaat | gtggtccacg | tgtatgaact | gctgctggag | 1320 |
| 230 | atgctgaatg | cccacgttgt | tgcctgggtg | aagtcctcca | tcaagggggtc | cagatgcagc | 1380 |
| 240 | ccggcagagg | acagtaaaag | caagaggggc | tcccagaacc | cacagttcca | gtga | 1434 |

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Input Set : A:\2355-124 Seq List.txt

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77 <210> SEQ ID NO: 2

78 <211> LENGTH: 1151

79 <212> TYPE: DNA

80 <213> ORGANISM: Homo sapiens

81 <400> SEQUENCE: 2

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83 atgaattaca gcaatccacag caatgtcaact aacttggaa gttgggcoctg tgggagagacc 60
85 acaaacccca atgtgttggg gccaacacct gggaaccttt ctcttttagt ggtccatcgc 120
87 caattatcac atctgtatgc ggaacctcaa aagagtcctt ggtgtgaagc aagatcgcta 180
89 caatccacct taactgtaaa cagagagaca ctgaaaagga aggttagtgg gaacgttgc 240
91 gccacccctt ttactgttcc aggttcaaa ggggatgtct actctgcgc tctctgcgc 300
93 gattacgaat ttcattatca ctatggagtc tggctgtgtg aaggatgtaa ggcctttttt 360
95 aaaaagagca ttccaggaca taatgattat attgtccag ctacaaatca gtgtacaatc 420
97 gataaaaccc ggcacagag ctgcacaggc tgcgcacttc ggaagtgtta cgaagtggga 480
99 atgttgaagt gtgtctcccg gagagagaga tctgggtacc gcttctgcg gagacagaga 540
101 cctcccgagg agagctgca ctgtgcgggc aaggccaaga gaagtggcgg ccacggcgcc 600
103 caattggcgg agtgtgtgt ggagggcctg agccccgagc agctagtgt caccctcctg 660
105 gagctgagc ccgaccatgt gctgacacgc cggccacgtg cggccctcac cagggcctcc 720
107 atgatgatct ccttgaccaa gttggccgac aaggagtgtg tacacatgat cagctgggac 780
109 acaagatttc caggtttgt ggagctcagc ctgttcagac aagtgcggtt ctctggagagc 840
111 tcttgatagg aattgttaat gatggggctg atgtggcgt caattgaaca ccccgccaag 900
113 ctactcttgg ctccagatct tgttctggac agggatgagg gaaaatgctt agaaggaatt 960
115 ctggaaat ct tccatgtgt cctggcaact acttcaaggt ttogagagtt aaaactccaa 1020
117 caaagaaat atctctgtgt caaggccatg atctgtctca attccagtat gtaacctctg 1080
119 ctccagagca ccaggcatgc tgacagcagc cgggaagctg ctcaactgtt gaaagcctg 1140
121 cccatgctt tgggttgggt gattggcaag agcgcatct cctccacaga gcaatccatg 1200
123 cgcctggcta cctccctgat gctcctgtcc caagtccagc atgcgaggtg a 1251

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125 <210> SEQ ID NO: 3

126 <211> LENGTH: 66

127 <212> TYPE: PRT

128 <213> ORGANISM: Homo sapiens

129 <400> SEQUENCE: 3

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131 Cys Ala Val Cys Ser Asp Tyr Ala Ser Gly Tyr His Tyr Gly Val Trp
132 1 5 10 15
134 Ser Cys Glu Gly Cys Lys Ala Phe Phe Lys Arg Ser Ile Gln Gly His
135 20 25 30
137 Asn Asp Tyr Ile Cys Pro Ala Thr Asn Gln Cys Thr Ile Asp Lys Asn
138 35 40 45
140 Arg Arg Lys Ser Cys Gln Ala Cys Arg Leu Arg Lys Cys Tyr Glu Val
141 50 55 60
143 Gly Met
144 65

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145 <210> SEQ ID NO: 4

146 <211> LENGTH: 233

147 <212> TYPE: PRT

148 <213> ORGANISM: Homo sapiens

149 <400> SEQUENCE: 4

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151 Leu Val Leu Thr Leu Leu Glu Ala Glu Pro Pro His Val Leu Ile Ser
152 1 5 10 15
154 Arg Pro Ser Ala Pro Phe Thr Glu Ala Ser Met Met Met Ser Leu Thr

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150          20          25          30
151 Lys Leu Ala Asp Lys Glu Leu Val His Met Ile Ser Trp Ala Lys Lys
152          35          40          45
153 Ile Pro Gly Phe Val Glu Leu Ser Leu Phe Asp Gln Val Arg Leu Leu
154          50          55          60
155 Glu Ser Cys Trp Met Glu Val Leu Met Met Gly Leu Met Trp Arg Ser
156 65          70          75          80
157 Ile Asp His Pro Gly Lys Leu Ile Phe Ala Pro Asp Leu Val Leu Asp
158          85          90          95
159 Arg Asp Glu Gly Lys Cys Val Glu Gly Ile Leu Glu Ile Phe Asp Met
160          100          105          110
161 Leu Leu Ala Thr Thr Ser Arg Phe Arg Glu Leu Lys Leu Gln His Lys
162          115          120          125
163 Glu Tyr Leu Cys Val Lys Ala Met Ile Leu Leu Asn Ser Ser Met Tyr
164          130          135          140
165 Pro Leu Val Thr Ala Thr Gln Asp Ala Asp Ser Ser Arg Lys Leu Ala
166          145          150          155          160
167 His Leu Leu Asn Ala Val Thr Asp Ala Leu Val Trp Val Ile Ala Lys
168          165          170          175
169 Ser Gly Ile Ser Ser Gln Gln Gln Ser Met Arg Leu Ala Asn Leu Leu
170          180          185          190
171 Met Leu Leu Ser His Val Arg His Ala Ser Asn Lys Gly Met Glu His
172          195          200          205
173 Leu Leu Asn Met Lys Cys Lys Asn Val Val Pro Val Tyr Asp Leu Leu
174          210          215          220
175 Leu Glu Met Leu Asn Ala His Val Leu
176          225          230
177 -2108- SEQ ID NO: 5
178 -2110- LENGTH: 477
179 -2112- TYPE: PRT
180 -2113- ORGANISM: Homo sapiens
181 -2400- SEQUENCE: 5
182 Met Asn Tyr Ser Ile Pro Ser Asn Val Thr Asn Leu Glu Gly Gly Pro
183          5          10          15
184 Gly Arg Gln Thr Thr Ser Pro Asn Val Leu Trp Pro Thr Pro Gly His
185          20          25          30
186 Leu Ser Pro Leu Val Val His Arg Gln Leu Ser His Leu Tyr Ala Glu
187          35          40          45
188 Pro Gln Lys Ser Pro Trp Cys Glu Ala Arg Ser Leu Glu His Thr Leu
189          50          55          60
190 Pro Val Asn Arg Glu Thr Leu Lys Arg Lys Val Ser Gly Asn Arg Cys
191 65          70          75          80
192 Ala Ser Pro Val Thr Gly Pro Gly Ser Lys Arg Asp Ala His Phe Cys
193          85          90          95
194 Ala Val Cys Ser Asp Tyr Ala Ser Gly Tyr His Tyr Gly Val Trp Ser
195          100          105          110
196 Cys Glu Gly Cys Lys Ala Phe Phe Lys Arg Ser Ile Gln Gly His Asn
197          115          120          125
198 Asp Tyr Ile Cys Pro Ala Thr Asn Gln Cys Thr Ile Asp Lys Asn Arg

```

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132      133      135      140
130 Arg Lys Ser Cys Gln Ala Cys Arg Leu Arg Lys Cys Tyr Glu Val Gly
141 145      150      155      160
131 Met Val Lys Cys Gly Ser Arg Arg Glu Arg Cys Gly Tyr Arg Leu Val
134      165      170      175
136 Arg Arg Gln Arg Ser Ala Asp Glu Gln Leu His Cys Ala Gly Lys Ala
137      180      185      190
138 Lys Arg Ser Gly Gly His Ala Pro Arg Val Arg Glu Leu Leu Asp
140      195      200      205
141 Ala Leu Ser Pro Gln Gln Leu Val Leu Thr Leu Leu Glu Ala Glu Pro
143      210      215      220
145 Pro His Val Leu Ile Ser Arg Pro Ser Ala Pro Phe Thr Glu Ala Ser
146 225      230      235      240
147 Met Met Met Ser Leu Thr Lys Leu Ala Asp Lys Glu Leu Val His Met
149      245      250      255
151 Ile Ser Trp Ala Lys Lys Ile Pro Gly Phe Val Glu Leu Ser Leu Phe
152      260      265      270
154 Asp Gln Val Arg Leu Leu Glu Ser Cys Trp Met Glu Val Leu Met Met
155      275      280      285
157 Gly Leu Met Trp Arg Ser Ile Asp His Pro Gly Lys Leu Ile Phe Ala
158      290      295      300
160 Pro Asp Leu Val Leu Asp Arg Asp Glu Gly Lys Cys Val Glu Gly Ile
161 305      310      315      320
163 Leu Glu Ile Phe Asp Met Leu Leu Ala Thr Thr Ser Arg Phe Arg Glu
164      325      330      335
166 Leu Lys Leu Gln His Lys Glu Tyr Leu Cys Val Lys Ala Met Ile Leu
167      340      345      350
169 Leu Asn Ser Ser Met Tyr Pro Leu Val Thr Ala Thr Gln Asp Ala Asp
170      355      360      365
172 Ser Ser Arg Lys Leu Ala His Leu Leu Asn Ala Val Thr Asp Ala Leu
173      370      375      380
175 Val Trp Val Ile Ala Lys Ser Gly Ile Ser Ser Gln Gln Gln Ser Met
176 385      390      395      400
178 Arg Leu Ala Asn Leu Met Leu Leu Ser His Val Arg His Ala Ser
179      405      410      415
181 Asn Lys Gly Met Glu His Leu Leu Asn Met Lys Cys Lys Asn Val Val
182      420      425      430
184 Pro Val Tyr Asp Leu Leu Leu Glu Met Leu Asn Ala His Val Leu Arg
185      435      440      445
187 Gly Cys Lys Ser Ser Ile Thr Gly Ser Glu Cys Ser Pro Ala Glu Asp
188      450      455      460
190 Ser Lys Ser Lys Glu Gly Ser Gln Asn Pro Gln Ser Gln
191 465      470      475

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193 <110> SEQ ID NO: 6

194 <111> LENGTH: 416

195 <112> TYPE: PRT

196 <113> ORGANISM: Homo sapiens

198 <400> SEQUENCE: 6

199 Met Asn Tyr Ser Ile Pro Ser Asn Val Thr Asn Leu Glu Gly Gly Pro

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300 1          5          10          15
301 Gly Arg Gln Thr Thr Ser Pro Asn Val Leu Trp Pro Thr Pro Gly His
302          20          25          30
303 Leu Ser Pro Leu Val Val His Arg Gln Leu Ser His Leu Tyr Ala Glu
304          35          40          45
305 Pro Gln Lys Ser Pro Trp Cys Glu Ala Arg Ser Leu Glu His Thr Leu
306          50          55          60
307 Pro Val Asn Arg Glu Thr Leu Lys Arg Lys Val Ser Gly Asn Arg Cys
308          65          70          75          80
309 Ala Ser Pro Val Thr Gly Pro Gly Ser Lys Arg Asp Ala His Phe Cys
310          85          90          95
311 Ala Val Cys Ser Asp Tyr Ala Ser Gly Tyr His Tyr Gly Val Trp Ser
312          100          105          110
313 Cys Glu Gly Cys Lys Ala Phe Phe Lys Arg Ser Ile Gln Gly His Asn
314          115          120          125
315 Asp Tyr Ile Cys Pro Ala Thr Asn Gln Cys Thr Ile Asp Lys Asn Arg
316          130          135          140
317 Arg Lys Ser Cys Gln Ala Cys Arg Leu Arg Lys Cys Tyr Glu Val Gly
318          145          150          155          160
319 Met Val Lys Cys Gly Ser Arg Arg Glu Arg Cys Gly Tyr Arg Leu Val
320          165          170          175
321 Arg Arg Gln Arg Ser Ala Asp Glu Gln Leu His Cys Ala Gly Lys Ala
322          180          185          190
323 Lys Arg Ser Gly Gly His Ala Pro Arg Val Arg Glu Leu Leu Leu Asp
324          195          200          205
325 Ala Leu Ser Pro Gln Gln Leu Val Leu Thr Leu Leu Glu Ala Glu Pro
326          210          215          220
327 Pro His Val Leu Ile Ser Arg Pro Ser Ala Pro Phe Thr Glu Ala Ser
328          225          230          235          240
329 Met Met Met Ser Leu Thr Lys Leu Ala Asp Lys Glu Leu Val His Met
330          245          250          255
331 Ile Ser Trp Ala Lys Lys Ile Pro Gly Phe Val Glu Leu Ser Leu Phe
332          260          265          270
333 Asp Gln Val Arg Leu Leu Glu Ser Cys Trp Met Glu Val Leu Met Met
334          275          280          285
335 Gly Leu Met Trp Arg Ser Ile Asp His Pro Gly Lys Leu Ile Phe Ala
336          290          295          300
337 Pro Asp Leu Val Leu Asp Arg Asp Glu Gly Lys Cys Val Glu Gly Ile
338          305          310          315          320
339 Leu Glu Ile Phe Asp Met Leu Leu Ala Thr Thr Ser Arg Phe Arg Glu
340          325          330          335
341 Leu Lys Leu Gln His Lys Glu Tyr Leu Cys Val Lys Ala Met Ile Leu
342          340          345          350
343 Leu Asn Ser Ser Met Tyr Pro Leu Val Thr Ala Thr Gln Asp Ala Asp
344          355          360          365
345 Ser Ser Arg Lys Leu Ala His Leu Leu Asn Ala Val Thr Asp Ala Leu
346          370          375          380
347 Val Trp Val Ile Ala Lys Ser Gly Ile Ser Ser Gln Gln Gln Ser Met
348          385          390          395          400

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RAW SEQUENCE LISTING ERROR SUMMARY
PATENT APPLICATION: US/08/826,361B

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Input Set : A:\2355-124 Seq List.txt
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Please Note:

Use of n and/or Xaa have been detected in the Sequence Listing. Please review the Sequence Listing to ensure that a corresponding explanation is presented in the <220> to <223> fields of each sequence which presents at least one n or Xaa.

Seq#:7; N Pos. 3,15,18
Seq#:8; N Pos. 12,18,24,27

VERIFICATION SUMMARY

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Input Set : A:\2355-124 Seq List.txt

Output Set: N:\CRF4\08212003\H826361B.raw

L:10 M:270 C: Current Application Number differs, Replaced Current Application No
L:10 M:271 C: Current Filing Date differs, Replaced Current Filing Date
L:388 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:7 after pos.:0
L:401 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:8 after pos.:0